

OIPE

RAW SEQUENCE LISTING

DATE: 07/31/2001

PATENT APPLICATION: US/09/835,992A

TIME: 13:14:40

Input Set : A:\L0461-7112.txt

Output Set: N:\CRF3\07312001\I835992A.raw

2 <110> APPLICANT: Obata, Yuichi
4 <120> TITLE OF INVENTION: ISOLATED NUCLEIC ACID MOLECULES ASSOCIATED WITH GASTRIC
CANCER AND
5 METHODS FOR DIAGNOSING AND TREATING GASTRIC CANCER
7 <130> FILE REFERENCE: L0461/7112
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/835,992A
C--> 11 <141> CURRENT FILING DATE: 2001-04-16
13 <150> PRIOR APPLICATION NUMBER: US 08/896,164
14 <151> PRIOR FILING DATE: 1997-07-17
16 <160> NUMBER OF SEQ ID NOS: 87
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 707
21 <212> TYPE: DNA
22 <213> ORGANISM: Homo sapiens
24 <220> FEATURE:
25 <221> NAME/KEY: Unsure
26 <222> LOCATION: (107)..(107)
27 <223> OTHER INFORMATION: n = a, c, g or t
29 <220> FEATURE:
30 <221> NAME/KEY: Unsure
31 <222> LOCATION: (154)..(154)
32 <223> OTHER INFORMATION: n = a, c, g or t
34 <220> FEATURE:
35 <221> NAME/KEY: Unsure
36 <222> LOCATION: (196)..(196)
37 <223> OTHER INFORMATION: n = a, c, g or t
39 <220> FEATURE:
40 <221> NAME/KEY: Unsure
41 <222> LOCATION: (215)..(215)
42 <223> OTHER INFORMATION: n = a, c, g or t
44 <220> FEATURE:
45 <221> NAME/KEY: Unsure
46 <222> LOCATION: (309)..(309)
47 <223> OTHER INFORMATION: n = a, c, g or t
49 <220> FEATURE:
50 <221> NAME/KEY: Unsure
51 <222> LOCATION: (453)..(453)
52 <223> OTHER INFORMATION: n = a, c, g or t
54 <220> FEATURE:
55 <221> NAME/KEY: Unsure
56 <222> LOCATION: (512)..(512)
57 <223> OTHER INFORMATION: n = a, c, g or t
59 <220> FEATURE:
60 <221> NAME/KEY: Unsure
61 <222> LOCATION: (543)..(543)
62 <223> OTHER INFORMATION: n = a, c, g or t
64 <220> FEATURE:

ENTERED

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Input Set : A:\L0461-7112.txt

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65 <221> NAME/KEY: Unsure
 66 <222> LOCATION: (550)..(550)
 67 <223> OTHER INFORMATION: n = a, c, g or t
 69 <220> FEATURE:
 70 <221> NAME/KEY: Unsure
 71 <222> LOCATION: (579)..(579)
 72 <223> OTHER INFORMATION: n = a, c, g or t
 74 <220> FEATURE:
 75 <221> NAME/KEY: Unsure
 76 <222> LOCATION: (588)..(588)
 77 <223> OTHER INFORMATION: n = a, c, g or t
 79 <220> FEATURE:
 80 <221> NAME/KEY: Unsure
 81 <222> LOCATION: (610)..(610)
 82 <223> OTHER INFORMATION: n = a, c, g or t
 84 <220> FEATURE:
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 87 <223> OTHER INFORMATION: n = a, c, g or t
 89 <220> FEATURE:
 90 <221> NAME/KEY: Unsure
 91 <222> LOCATION: (650)..(650)
 92 <223> OTHER INFORMATION: n = a, c, g or t
 94 <220> FEATURE:
 95 <221> NAME/KEY: Unsure
 96 <222> LOCATION: (654)..(654)
 97 <223> OTHER INFORMATION: n = a, c, g or t
 99 <220> FEATURE:
 100 <221> NAME/KEY: Unsure
 101 <222> LOCATION: (671)..(671)
 102 <223> OTHER INFORMATION: n = a, c, g or t
 104 <400> SEQUENCE: 1
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 W--> 106 gggaagctat gcgaaattat ttaaaagagc gaggggatca aacagtnctt attcttcatg 120
 W--> 107 caaaagttgc acagaagtca tatggaaatg aaanaagggt tttttgccca cctccttggtg 180
 W--> 108 tatatcttat gggcantgga tggaagaaaa aaaangaaca aatggaacgc gatggttggt 240
 109 ctgaacaaaa gtctcaaccg tgtgcattta ttgggatagg aaatagtgac caaaaaatgc 300
 W--> 110 agcagctana cttggaagga aagaactatt gcacagccaa aacattgtat atatctgact 360
 111 cagacaagcg aaagcacttc atgttgctctg taaagatggt ctatggcaac agtgatgaca 420
 W--> 112 ttggtgtggt cctcagcaag cgaataaaag tcntctccaa accttccaaa aagaacagtc 480
 W--> 113 attgaaaaat gctgacttat gcattgcctc angaacaaag gtggctctgt ttaatcgact 540
 W--> 114 acnatcccan acagttagta ccagatactt gcatgttana aggaggtnat tttcatgcca 600
 W--> 115 gttcacagcn gtggggagagc ttttttatct anctcttgga tgatgatgan tccnaaggag 660
 W--> 116 aagaattcac ngtccgagat ggctacatcc attatggaca aacagtc 707
 119 <210> SEQ ID NO: 2
 120 <211> LENGTH: 551
 121 <212> TYPE: DNA
 122 <213> ORGANISM: Homo sapiens
 124 <220> FEATURE:

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Input Set : A:\L0461-7112.txt

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125 <221> NAME/KEY: Unsure
 126 <222> LOCATION: (200)..(200) ✓
 127 <223> OTHER INFORMATION: n = a, c, g or t
 129 <220> FEATURE:
 130 <221> NAME/KEY: Unsure
 131 <222> LOCATION: (352)..(352) ✓
 132 <223> OTHER INFORMATION: n = a, c, g or t
 134 <220> FEATURE:
 135 <221> NAME/KEY: Unsure
 136 <222> LOCATION: (455)..(455)
 137 <223> OTHER INFORMATION: n = a, c, g or t ✓
 139 <220> FEATURE:
 140 <221> NAME/KEY: Unsure
 141 <222> LOCATION: (544)..(544) ✓
 142 <223> OTHER INFORMATION: n = a, c, g or t
 144 <220> FEATURE:
 145 <221> NAME/KEY: Unsure
 146 <222> LOCATION: (546)..(546) ✓
 147 <223> OTHER INFORMATION: n = a, c, g or t
 149 <400> SEQUENCE: 2
 150 actgtggctt ctgcatttca aatcagcact tgcagggaga caacgggggtt tttgaatagt 60
 151 atcacctggt atgaaaagtt ttcccaagaa accacaaacg attgttcatt ttttctcctt 120
 152 ttttggttaac tttttgccac actcaagtca gtttaagtcc tagcaaaaag acggtagtta 180
 W--> 153 ggataccact gtggctgtan atgatgtgac actggttgaa tttgtgctgg cgtttgtgta 240
 154 acttcctcgc ctgtttgtgt ttgattcgtt aggggggcacc tggcttgaat tggctcgaag 300
 W--> 155 gattgctcct gctgcactgc aatgtggccg cggccctggt tctggtgtgt angtaaaggt 360
 156 aaggctggtg gaataaatga ttccaccatt tcggaccaa gttactggaa cctggactgg 420
 W--> 157 ttgccggacc catctccaac cttctcggaa tgcanaaatg tctgggacga cacagaacat 480
 158 acctctccac acctgtacat aatttcagct tctacatccc caaaccacac tcgtaaattt 540
 W--> 159 ggantnaaaa t 551
 162 <210> SEQ ID NO: 3
 163 <211> LENGTH: 478
 164 <212> TYPE: DNA
 165 <213> ORGANISM: Homo sapiens
 167 <220> FEATURE:
 168 <221> NAME/KEY: Unsure
 169 <222> LOCATION: (39)..(39) ✓
 170 <223> OTHER INFORMATION: n = a, c, g or t
 172 <220> FEATURE:
 173 <221> NAME/KEY: Unsure
 174 <222> LOCATION: (100)..(100) ✓
 175 <223> OTHER INFORMATION: n = a, c, g or t
 177 <220> FEATURE:
 178 <221> NAME/KEY: Unsure
 179 <222> LOCATION: (135)..(135) ✓
 180 <223> OTHER INFORMATION: n = a, c, g or t
 182 <220> FEATURE:
 183 <221> NAME/KEY: Unsure
 184 <222> LOCATION: (147)..(147) ✓

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185 <223> OTHER INFORMATION: n = a, c, g or t
187 <220> FEATURE:
188 <221> NAME/KEY: Unsure ✓
189 <222> LOCATION: (170)..(170)
190 <223> OTHER INFORMATION: n = a, c, g or t
192 <220> FEATURE:
193 <221> NAME/KEY: Unsure ✓
194 <222> LOCATION: (206)..(206)
195 <223> OTHER INFORMATION: n = a, c, g or t
197 <220> FEATURE:
198 <221> NAME/KEY: Unsure ✓
199 <222> LOCATION: (270)..(270)
200 <223> OTHER INFORMATION: n = a, c, g or t
202 <220> FEATURE:
203 <221> NAME/KEY: Unsure ✓
204 <222> LOCATION: (339)..(339)
205 <223> OTHER INFORMATION: n = a, c, g or t
207 <220> FEATURE:
208 <221> NAME/KEY: Unsure ✓
209 <222> LOCATION: (347)..(347)
210 <223> OTHER INFORMATION: n = a, c, g or t
212 <220> FEATURE:
213 <221> NAME/KEY: Unsure ✓
214 <222> LOCATION: (357)..(357)
215 <223> OTHER INFORMATION: n = a, c, g or t
217 <220> FEATURE:
218 <221> NAME/KEY: Unsure ✓
219 <222> LOCATION: (360)..(360)
220 <223> OTHER INFORMATION: n = a, c, g or t
222 <220> FEATURE:
223 <221> NAME/KEY: Unsure ✓
224 <222> LOCATION: (364)..(364)
225 <223> OTHER INFORMATION: n = a, c, g or t
227 <220> FEATURE:
228 <221> NAME/KEY: Unsure ✓
229 <222> LOCATION: (389)..(389)
230 <223> OTHER INFORMATION: n = a, c, g or t
232 <220> FEATURE:
233 <221> NAME/KEY: Unsure ✓
234 <222> LOCATION: (395)..(395)
235 <223> OTHER INFORMATION: n = a, c, g or t
237 <220> FEATURE:
238 <221> NAME/KEY: Unsure ✓
239 <222> LOCATION: (417)..(418)
240 <223> OTHER INFORMATION: n = a, c, g or t
242 <220> FEATURE:
243 <221> NAME/KEY: Unsure ✓
244 <222> LOCATION: (422)..(422)
245 <223> OTHER INFORMATION: n = a, c, g or t

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247 <220> FEATURE:
248 <221> NAME/KEY: Unsure
249 <222> LOCATION: (424)..(424)
250 <223> OTHER INFORMATION: n = a, c, g or t
252 <220> FEATURE:
253 <221> NAME/KEY: Unsure
254 <222> LOCATION: (442)..(442)
255 <223> OTHER INFORMATION: n = a, c, g or t
257 <220> FEATURE:
258 <221> NAME/KEY: Unsure
259 <222> LOCATION: (451)..(451)
260 <223> OTHER INFORMATION: n = a, c, g or t
262 <220> FEATURE:
263 <221> NAME/KEY: Unsure
264 <222> LOCATION: (463)..(463)
265 <223> OTHER INFORMATION: n = a, c, g or t
267 <220> FEATURE:
268 <221> NAME/KEY: Unsure
269 <222> LOCATION: (468)..(468)
270 <223> OTHER INFORMATION: n = a, c, g or t
272 <400> SEQUENCE: 3
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W--> 274 atcacctggt atgaaaagtt tteccaagaa accacaaacn attgttcatt ttttctcctt 120
W--> 275 ttttggttaac ttttngccac actcaantca gtttaagtcc tagcaaaaan acggtagtta 180
W--> 276 ggataccact gtggctgtaa atgatntgac actgggtgaa tttgtgctgg cgtttgtgta 240
W--> 277 acttccctcg ctgtttgtgt ttgattcgtt aggggggcacc tggcttgaat tggctcgaag 300
W--> 278 gattgctcct gctgcactgc aatgtggcgc cgggcctgnt tcttatntgt tgtaaangtn 360
W--> 279 aggntgggtgg aataaatgat tccatcatnt cggancgaag ttgctgggaa ctggganngg 420
W--> 280 tngncggaac catctccgac cncccggaaa ngcagaagtg ttngtggnag accggaac 478
283 <210> SEQ ID NO: 4
284 <211> LENGTH: 707
285 <212> TYPE: DNA
286 <213> ORGANISM: Homo sapiens
288 <220> FEATURE:
289 <221> NAME/KEY: Unsure
290 <222> LOCATION: (88)..(88)
291 <223> OTHER INFORMATION: n = a, c, g or t
293 <220> FEATURE:
294 <221> NAME/KEY: Unsure
295 <222> LOCATION: (147)..(147)
296 <223> OTHER INFORMATION: n = a, c, g or t
298 <220> FEATURE:
299 <221> NAME/KEY: Unsure
300 <222> LOCATION: (157)..(157)
301 <223> OTHER INFORMATION: n = a, c, g or t
303 <220> FEATURE:
304 <221> NAME/KEY: Unsure
305 <222> LOCATION: (163)..(163)
306 <223> OTHER INFORMATION: n = a, c, g or t

```

→ Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

VERIFICATION SUMMARY

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Input Set : A:\L0461-7112.txt

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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1

L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:275 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:276 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3

L:360 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

L:361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

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L:367 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

L:368 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

L:369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

L:370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:691 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:693 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:694 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:695 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:696 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:697 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5

L:822 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:823 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:824 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:825 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:826 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:828 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:830 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:831 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

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L:868 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:870 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:871 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:927 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8